

REPLACEMENT SHEET

1/12

1 AAGCTTCGGG GCCTCTGCAG AKAAGTTGTG CAAACACTTC CTCTCTGGAC
 51 AAGGAGGAAT GCAGGAGGCC ACCGCGTGCA GTACATCTTG GAGTGTGGGA
 101 GGGATGTGCC TGCACTTGTG AACGGGGGCC AGGAGAATGA GGGCCCAACC
 151 AAGCCCGGCA GTGCCAGTA GATGCAGAGA GGCACCCGTG CCCCAGTGA
 201 GGGCACAGCC AGTGGGCATC CCTGAGGCCC AGGGGCCCGT TCCTCTCCAC
 251 TCCCAACAGA TGCGTCTGCT GTGGGGCCCT GGGAGGCCCT TCCTCGCCCT
 301 GCTTCTGCTG GTTCCATCA AGCAAGTTAC AGGATCGCTC CTCAGGAGA
 351 CAAGTCAGAA GTGGGCTAAT TATAAGGAGA AGTGTCTGGA AGACTTGCAC
 401 AATAGACTTT CTGGCATAIT TTGTAATGGG ACATTGTATC GGTATGTGTG
 451 CTGGCCTCAT TCTTATCCTG GAAATGTCTC TGTTCCTGTG CCTTCATCT
 501 TACCTTGGTG GAATGCAGAG AGCCGAGGAA GGGCCTACAG ACATGTCTTG
 551 GCTCAGGGGA CTGGCGAGAC GCGAGAGAAC ACCACAGATA TTGGCAGGA
 601 TGAATCAGAA TGCTCAGAGA ACCACAGCTT CAGACAAAAC GTGGATCACT
 651 ACGCCTTGCT ATACACCTTG CAGCTGATGT ACATGTGGGG CTACTCCGTG
 701 TCTCTCATCT CCGTCTTCTT GGCTCTTACA CTCTCTTGT TCCTTCGAAA
 751 ACTGCATTGC ACACGCAATT ACATCCACAT GAACCTGTTC GCTTCGTCA
 801 TCCTGAAGT TCTGGCTGTC CTGGTGAAGG ACATGGTCTC CCACACTCT
 851 TACTCCAAGA GGCOCGATGA TGAGAGTGA TGGATGTCAT ATCTGTCAGA
 901 GACATCCGTC TCCTGTGCGT CCGTCCAGGT CCTCTGCAC TACTTTGTGG
 951 GCACCAATCA CTTGTGGCTG CTGGTTGAAG GACTTTACCT CCACACTCTG
 1001 CTGGAGCCCA CAGTGTTCCT TGAAGGCGG CTGTGGCCCA AGTACCTGGT
 1051 GGTGGGTGGG GCGTCCCCCA TGCTGTTTGT TATTCCTGG GGTTTGGCCC
 1101 GTGCACACT GGAGAACACA CGGTGCTGGG CCACAAATGG GAACGTGAAA
 1151 ATCTGTGGGA TCATCAGAGG ACCCATGCTG CTTTGGTAA CAGTTAATT
 1201 CTTCATCTTC CTCAGATTG TCAAGCTTCT CATTCTAAG CTCAGAGCTC
 1251 ATCAGATGTG CTCAGAGAC TACAAATACA GATTGGCGAA ATCAAAGTTG
 1301 CTCCTCATTC CTTGTGGGG GGTTCATGAG GTCCTCTTCA CTTCTTCCC
 1351 CGACGACCAA GTTCAAGGAT TTTCAAAACG TATTCGACTC TTCAATCAGC
 1401 TGACACTGAG CTCTGTCCAC GGATTTCTGG TGGCCTTGCA GTATGGCTTT

FIGURE 1